The reduction of biological production induced by mesoscale mixing: a modelling study in the Benguela upwelling.

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Abstract

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Recent studies, both based on remote sensed data and coupled models, showed a re-12 duction of biological productivity due to vigorous horizontal mixing in upwelling systems. In order to better understand this phenomenon, we have considered a system of oceanic 14 flow in the Benguela area coupled with a simple biogeochemical model of Nutrient-Phyto-15 Zooplankton (NPZ) type. For the flow three different surface velocity fields are considered: 16 one derived from satellite altimetry data, and the other two from a regional numerical 17 model at two different spatial resolutions. We computed horizontal particle dispersion in terms of Lyapunov Exponents, and analyzed their correlations with phytoplankton 19 concentrations. Our modelling approach confirms that in the south Benguela, there is a reduction of biological activity when stirring is increased. Two-dimensional offshore advection seems to be the dominant process involved. In the northern area, other factors not taken into account in our simulation are influencing the ecosystem. We provide explanations for these results in the context of studies performed in other Eastern Boundary upwelling areas.

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e 1. Introduction

Marine ecosystems of the Eastern Boundary Upwelling zones are well known for their major contribution to the world ocean productivity. They are characterized by winddriven upwelling along the coast of cold nutrient-rich waters that supports elevated plankton and pelagic fish production (Mackas et al., 2006). Variability is introduced by strong
advection along the shore, physical forcings by local and large scales winds, and high suband mesoscale activities over the continental shelf and beyond, linking the coastal domain
with the open ocean.

The Benguela Upwelling System (BUS) is one of the four major Eastern Boundary
Systems (EBUS) of the world. The coastal area of the Benguela ecosystem extends from
southern Angola (around 17°S) along the west coast of Namibia and South Africa (36°S).

Systems (EBUS) of the world. The coastal area of the Benguela ecosystem extends from southern Angola (around 17°S) along the west coast of Namibia and South Africa (36°S). It is surrounded by two warm temperate boundary currents, the Angola Current in the north, and the Agulhas Current in the south. The BUS can itself be subdivided into two subdomains by the powerful Luderitz upwelling cell (Hutchings et al., 2009). Most of the biogeochemical activity occurs within the upwelling front and the coast, although it can be extended further offshore toward the open ocean by the numerous filamental structures developing offshore (Monteiro, 2009). In the BUS, as in the other major upwelling areas, a high mesoscale activity due to eddies and filaments is observed and impacts strongly on marine planktonic ecosystem over the shelf and beyond (Brink and Cowles, 1991; Martin, 2003; Sandulescu et al., 2008; Rossi et al., 2009).

The purpose of this study is to analyze the impact of the horizontal stirring on the phytoplankton dynamics in the BUS. Recently, Rossi et al. (2008, 2009), using satellite data
of the ocean surface, suggested that mesoscale activity has a negative effect on chlorophyll
standing stocks in the EBUS. This was obtained by correlating remote sensed chlorophyll
data with a Lagrangian measurement of lateral stirring in the surface ocean (see Methods
section below). This result was unexpected since mesoscale transport, particularly due
to eddies, has been related to higher planktonic production and stocks in the open ocean

(McGillicuddy et al., 2007) as well as off a major EBUS (Correa-Ramirez et al., 2007). A more recent and thorough study performed by Gruber et al. (2011) in the California and the Canary current systems detailed the initial results from Rossi et al. (2008, 2009). 55 Based on satellite derived estimates of net Primary Production, of upwelling strength and of Eddy Kinetic Energy (EKE) as a measure the intensity of mesoscale activity, they confirmed the suppressive effect of mesoscale structures on biological production in upwelling areas. The mechanism behind this observation was investigated using 3D eddy resolving coupled models. The eddies tend to export offshore and downward a certain pool of nutrients not being effectively used by the biology in the coastal areas. This process they 61 called "nutrients leakage" is also having a negative feedback effect by diminishing the nutrients available in the deep waters being re-upwelled continuously. 63 In our work, we focused on the Benguela area, being the most contrasting area of 64 all EBUS in term of mixing intensity. Although mechanisms involved occur in the 3D

In our work, we focused on the Benguela area, being the most contrasting area of all EBUS in term of mixing intensity. Although mechanisms involved occur in the 3D space, the initial observation of this suppressive effect was based only on two-dimensional (2D) datasets (Rossi et al., 2008). Here we use 2D numerical analysis in a simple semi-realistic framework to test the effect of horizontal advection versus biological dynamics. Meanwhile, since vertical dimension is crucial in upwelling areas, it was introduced in our model in a simplified way by considering a source term with an intensity and spatial distribution corresponding to the upwelling characteristics. Indeed other theoretical studies in idealized 2D settings display also negative correlation between mixing and biomass (Tél et al., 2005; MacKiver and Neufeld, 2009). Contrarily to EKE which is an Eulerian diagnostic tool, we used here a Lagrangian measurement of mesoscale intensity. It has been demonstrated as a powerful tool to study patchy chlorophyll distributions due to dynamical structures at mesoscale, such as upwelling filaments (Calil and Richards, 2010). Different velocity fields were considered, one obtained from satellite and others from numerical simulations. The robustness of our results with respect to spatial resolution is tested by using two numerical velocity datasets at different resolution. Our results are

compared with real distributions of chlorophyll (a metric for phytoplankton) obtained from SeaWiFS satellite.

This paper is organized as follows. In Sec. 2 we describe the different data sets for our analysis. Sec. 3 contains the methodology, including the Finite-Size Lyapunov exponents, and the plankton numerical model. Then, in Sec. 4 our results are presented and discussed in the context of existing bibliography. Finally in Sec. 5, we summed-up our main findings.

2. Satellite and simulated data.

A total of three sources of two-dimensional velocity data sets in the surface of the 88 Benguela area were used: two were obtained from the numerical model ROMS (Regional Ocean Model System), and the other one from a combined satellite product. ROMS is 90 a free surface, hydrostatic, primitive equation model, and the run used here was eddy 91 resolving but climatologically forced (Gutknecht et al., 2011). At each grid point, linear 92 horizontal resolution is the same in both the longitudinal, ϕ , and latitudinal, θ , directions, which leads to angular resolutions $\Delta \phi = \Delta_0$ and $\Delta \theta = \Delta \phi \cos \theta$. The numerical model 94 was run onto 2 different grids: a coarse one at $\Delta_0 = 1/4^{\circ}$, and a finer one at $\Delta_0 = 1/12^{\circ}$ 95 of spatial resolution. In the following we label the data set from the coarser resolution 96 as ROMS1/4, and the finer one as ROMS1/12. In both of them, vertical resolution is 97 variable with 30 layers in total. Only data from the upper layer were used. The third set of velocity data are surface currents computed from a combination of wind-driven 99 Ekman currents, at 15 m depth, derived from Quickscat wind estimates, and geostrophic 100 currents calculated using time variable Sea Surface Heights (SSH) obtained from satellite 101 (Sudre and Morrow, 2008). These SSH were calculated from mapped altimetric sea level 102 anomalies combined with a mean dynamic topography. This velocity field, labeled as 103 Satellite 1/4, covers a period from June 2002 to June 2005 with a spatial resolution of 104 $\Delta_0 = 1/4^{\circ}$ in both longitudinal and latitudinal directions. 105

To validate simulated biological fields we used a three-year-long time series, from January 2002 to January 2005, of ocean color data. Phytoplankton pigment concentration
(chlorophyll-a) are obtained from monthly SeaWiFS (Sea viewing Wide Field-of-view Sensor) products, generated by the NASA Goddard Earth Science (GES)/Distributed Active
Archive Center (DAAC). Gridded global data were used with a resolution of approximately 9 by 9 km.

112 3. Methodology.

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3.1. Finite Size Lyapunov Exponents (FSLEs).

FSLEs (Artale et al., 1997; Aurell et al., 1997; Boffetta et al., 2001) provides a measure of dispersion, and thus of stirring and mixing, as a function of the spatial resolution, serving to isolate the different regimes corresponding to different length scales of the oceanic flows, as well as identifying the Lagrangian Coherent Structures (LCSs) present in the data. FSLE are computed from τ , the time required for two particles of fluid (one of them placed at \mathbf{x}) to separate from an initial (at time t) distance of δ_0 to a final distance of δ_f , as

$$\lambda(\mathbf{x}, t, \delta_0, \delta_f) = \frac{1}{\tau} \log \frac{\delta_f}{\delta_0}.$$
 (1)

It is natural to choose the initial points \mathbf{x} on the nodes of a grid with lattice spacing 114 coincident with the initial separation of fluid particles δ_0 . Then, values of λ are obtained 115 in a grid with lattice separation δ_0 . In this work we take always the resolution of the FSLE 116 field, δ_0 , equal to the resolution of the velocity field, Δ_0 . Other choices of parameter 117 are possible and δ_0 can take any value, even much smaller than the resolution of the 118 velocity field (Hernández-Carrasco et al., 2011a). This opens many possibilities that will 119 not be explored in this work, since we focus here in the primary production, and, in 120 some instances, the influence of the data resolution, not on the resolution of the FSLEs 121 computation. 122

The field of FSLEs thus depends on the choice of two length scales: the initial, δ_0 123 and the final δ_f separations. As in previous works (d'Ovidio et al., 2004, 2009; Rossi 124 et al., 2008; Hernández-Carrasco et al., 2011a) we will focus on transport processes at 125 mesoscale, so that δ_f is taken as about 110 km, which is the order of the size of mesoscale 126 eddies at mid latitudes. To compute λ we need to know the trajectories of the particles 127 which gives Lagrangian character to this quantity. The equations of motion that describe 128 the horizontal evolution of particle trajectories in longitudinal and latitudinal spherical 129 coordinates, $\mathbf{x} = (\phi, \lambda)$, are: 130

$$\frac{d\phi}{dt} = \frac{u(\phi, \theta, t)}{R\cos\theta},\tag{2}$$

$$\frac{d\phi}{dt} = \frac{u(\phi, \theta, t)}{R\cos\theta}, \qquad (2)$$

$$\frac{d\theta}{dt} = \frac{v(\phi, \theta, t)}{R}, \qquad (3)$$

where u and v represent the eastwards and northwards components of the surface velocity 131 field, and R is the radius of the Earth (6400 km). 132

The ridges of the FSLE field can be used to define the Lagrangian Coherent Struc-133 tures (LCSs) (Haller and Yuan, 2000; d'Ovidio et al., 2004, 2009; Tew Kai et al., 2009; 134 Hernández-Carrasco et al., 2011a), useful to characterize the flow from the Lagrangian 135 point of view (Joseph and Legras, 2002; Koh and Legras, 2002). In fact, since we are only 136 interested in the ridges with large values of FSLE, the ones which significantly affect mix-137 ing, LCSs can be obtained as the regions with high values of FSLE, which have a line-like 138 shape. We will compute FSLEs integrating backwards-in-time the particle trajectories, 139 since attracting LCSs associated to this (the unstable manifolds) have a direct physical in-140 terpretation (Joseph and Legras, 2002; d'Ovidio et al., 2004, 2009). Tracers (chlorophyll, temperature, ...) spread along the attracting LCSs, thus creating their typical filamental 142 structure (Lehan et al., 2007; Calil and Richards, 2010). 143

3.2. The Biological model 144

The plankton model is similar to the one used in previous studies by Oschlies and 145 Garçon (1998, 1999) and Sandulescu et al. (2007, 2008). It describes the interaction of a three-level trophic chain in the mixed layer of the ocean, including, phytoplankton P, zoo-plankton Z and dissolved inorganic nutrient N, whose concentrations evolve in time according to the following equations:

$$\frac{dN}{dt} = F_N = \Phi_N - \beta \frac{N}{\kappa_N + N} P + \mu_N \left((1 - \gamma) \frac{\alpha \eta P^2}{\alpha + n P^2} Z + \mu_P P + \mu_z Z^2 \right), \tag{4}$$

$$\frac{dP}{dt} = F_P = \beta \frac{N}{\kappa_N + N} P - \frac{\alpha \eta P^2}{\alpha + \eta P^2} Z - \mu_P P, \tag{5}$$

$$\frac{dZ}{dt} = F_Z = \gamma \frac{\alpha \eta P^2}{\alpha + \eta P^2} Z - \mu_Z Z^2. \tag{6}$$

where the dynamics of the nutrients, Eq. (4), is determined by nutrient supply due to the vertical mixing Φ_N , its uptake by phytoplankton (2nd term) and its recycling by bacteria from sinking particles (3rd term). Vertical mixing which brings nutrients from lower layers into the mixed surface layer of the ocean is parameterized in the model (see below), since the hydrodynamical part considers only horizontal 2D transport. Terms in Eq. (5) stand for phytoplankton growth by consuming N, the grazing by zooplankton, and its natural mortality. The last equation, Eq. (6), represents zooplankton growth by consuming phytoplankton minus its quadratic mortality.

A crucial part of this model comes in the vertical mixing, Φ_N , since it mimics the upwelling. Assuming constant nutrient concentration N_0 below the mixed layer, this term reads:

$$\Phi_N = S(\mathbf{x}, t)(N_0 - N),\tag{7}$$

where the temporally and spatially dependent (on the two dimension location \mathbf{x}) function S determines the strength and the horizontal spatial distribution of vertical mixing in the model, thus specifying the upwelling characteristics. Thus, the vertical dynamics is introduced in our two-dimensional model via this function S. Upwelling intensity along the coast is characterized by a number of cells of enhanced vertical ekman driven transport that are associated with similar fluctuations of the alongshore wind (Demarcq et al., 2003; Veitch et al., 2009). Following these results, we use a function S which is different from

zero in a strip 0.5° wide from the coast. Its spatial dependence along the coast is plotted in Fig. 1. For the temporal dependence, S alternates between the two configurations 166 displayed in Fig. 1 one for winter and another for summer. Six separate upwelling cells can 167 be discerned in the figure, with peaks at approximately 33°S, 31°S, 27.5°S, 24.5°S, 21.5°S, 168 17.5°S, which are known with the following names: Peninsula, Columbine+Namaqua, 169 Luderitz, Walvis Bay, Namibia and Cunene, respectively. Luderitz being the strongest. 170 The dynamical system given by Eqs. (4,5,6), for values of S in the range shown on 171 Fig. 1, evolves towards equilibrium for N, P and Z. But S is not fixed and its spatial 172 dependence introduces a coupling with the hydrodynamics. The transient time to reach 173 equilibrium is typically 60 days with the initial concentrations used (see Sec. 3.3). The 174 parameters are set following a study by Pasquero et al. (2004) and are listed in Table 1.

parameter	value
β	$0.66 \mathrm{day^{-1}}$
η	$1.0 \text{ (mmol N m}^{-3})^{-2} \text{ day}^{-1}$
γ	0.75
a	$2.0~\mathrm{day^{-1}}$
k_N	$0.5~\rm mmol~N~m^{-3}$
μ_N	0.2
μ_P	$0.03 \ \rm day^{-1}$
μ_Z	$0.2 \text{ (mmol N m}^{-3})^{-2} \text{ day}^{-1}$
N_0	$8.0~\mathrm{mmol~N~m^{-3}}$

Table 1: List of parameters used in the biological model.

176 3.3. Coupling hydrodynamical and biological model in Benguela.

The evolution of the concentrations within a flow is determined by the coupling between the hydrodynamical and biological models, and it is performed by the advection-

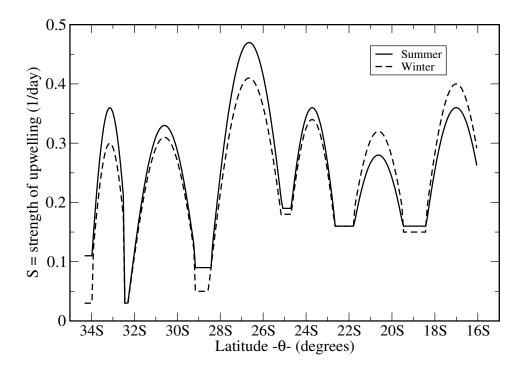


Figure 1: Shape and values of the strength (S) of the upwelling cells used in the simulations for winter and summer seasons (following Veitch et al. (2009)).

reaction-diffusion system. Thus, the complete model is given by the following system of 179 partial differential equations:

$$\frac{\partial N}{\partial t} + \mathbf{v}\nabla N = F_N + D\nabla^2 N,\tag{8}$$

$$\frac{\partial P}{\partial t} + \mathbf{v}\nabla P = F_P + D\nabla^2 P,\tag{9}$$

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$$\frac{\partial P}{\partial t} + \mathbf{v}\nabla P = F_P + D\nabla^2 P, \tag{9}$$

$$\frac{\partial Z}{\partial t} + \mathbf{v}\nabla Z = F_Z + D\nabla^2 Z. \tag{10}$$

The biological model is the one described before by the functions F_N , F_P and F_Z . 181 Horizontal advection is the 2D velocity \mathbf{v} , which is obtained from satellite data or from 182

the ROMS model. We add also an eddy diffusion term, via the ∇^2 operator, acting on N, P, and Z to incorporate the small-scale turbulence, which is not explicitly taken into account by the velocity fields used.

The eddy diffusion coefficient, D, is given by Okubo's formula (Okubo, 1971), $D(l) = 2.055 * 10^{-4} l^{1.15}$, where l is the value of the resolution, in meters, corresponding to the angular resolution $l = \Delta_0$. The formula gives the values $D=26.73 m^2/s$ for Satellite1/4 and ROMS1/4, and $D=7.4 m^2/s$ for ROMS1/12.

The coupled system Eqs. (8,9,10) is solved numerically by the semi-Lagrangian algo-190 rithm described in Sandulescu et al. (2007), combining Eulerian and Lagrangian schemes. 191 The initial concentrations of the tracers were taken from Koné et al. (2005) and they 192 are $N_0=1\ mmolNm^{-3}$, $P_0=0.1\ mmolNm^{-3}$, and $Z_0=0.06\ mmolNm^{-3}$. The in-193 flow conditions at the boundaries are specified in the following way: into the eastern, 194 western, and southern parts of the computation domain fluid parcels enter with very 195 poor biomasses concentration: $N_L = 0.01N_0 \ mmol Nm^{-3}$, $P_L = 0.01P_0 \ mmol Nm^{-3}$, and 196 $Z_L = 0.01 Z_0 \ mmol Nm^{-3}$. Across the northern boundary, fluid parcels enter with higher concentrations $N_H = 5 \ mmol Nm^{-3}$, $P_H = 0.1 \ mmol Nm^{-3}$, and $Z_H = 0.06 \ mmol Nm^{-3}$ 198 according with the values given by CARS for the Benguela system (Condie and Dunn, 190 2006). The integration time step is dt = 6 hours. 200

201 4. Results and discussion.

In this section we first compute the FSLEs on the velocity fields to quantify the horizontal stirring activity over the area. Then we analyze the results of the coupled biological-hydrodynamic model. Finally we investigate the relation between horizontal stirring activity and biological productivity.

206 4.1. Horizontal activity

We have computed the FSLE with a initial separation of particles equal to the spatial resolution of each velocity fields ($\delta_0 = 1/4^{\circ}$ for Satellite1/4 and ROMS1/4, and $\delta_0 = 1/12^{\circ}$

for ROMS1/12). As already mentioned, the final distance is always chosen to focus on transport processes by mesoscale structures at mid latitudes, $\delta_f = 1^{\circ}$. The areas of more 210 intense horizontal mixing can be identified using time averages of the backward FSLEs 211 (d'Ovidio et al., 2004). Figure 2 allows an easy characterization of sub-regions with dif-212 ferent horizontal mixing activity in the Benguela system. Areas of large average values of 213 FSLEs are identified as exhibiting an intense horizontal stirring or mesoscale activity. We 214 confirm the results of Rossi et al. (2009) by using different velocity data sets. Although 215 there are visible differences in the detailed patterns, good agreement between all datasets 216 is shown when computing the spatial correlation: for instance, correlation coefficient R^2 217 between FSLEs map from Satellite1/4 and from ROMS1/4 is 0.81. Correlation coeffi-218 cients between Satellite1/4 and ROMS1/12 on one hand, and between ROMS1/4 and 219 ROMS1/12 on the other hand, are lower (0.61 and 0.77 respectively) since the FSLE were 220 computed on a different resolution. More details on the effect on the grid resolution when 221 computing FSLEs can be found in Hernández-Carrasco et al. (2011a). For all data sets 222 high mixing values are observed in the southern region, while the northern area displays 223 significantly lower values. Note that the separation is well marked for Satellite 1/4 where 224 the line between the two areas is around 27°. In the case of the ROMS data sets, the 225 mixing activity is more homogeneously distributed, although the north-south gradient 226 is still present. We associate this difference with the injection of strong and numerous 227 Agulhas rings into the south of the area from the Agulhas retroflection.

The latitudinal behavior of mixing along the coastal upwelling can be seen in Fig. 3.

This was performed by computing the longitudinal averages of the plots in Fig. 2 for two coastally oriented strips, of 3° and 6° width, respectively. It is clear that horizontal mixing decreases as latitude decreases. Note that there are differences in the mixing values (FSLEs) depending on the type of data, their resolution and the grid size of FSLE computation. In general, considering velocities with the same resolution, the lower values correspond to Satellite 1/4 as compared to ROMS1/4. On average, values of mixing from

ROMS1/4 are larger than those from ROMS1/12, whereas we would expect the opposite considering the higher resolution of the latter simulation favouring small scales processes. However a caveat here is that FSLE were not computed on the same resolution, so there are not directly comparable. Note also that a low-mixing region is observed from 28° to 30°S on all calculations. It seems to indicate that the ROMS model is representing pretty well the spatial variability of the mixing. As proposed in a recent study by Titaud et al. (2011), these preliminary results indicate that FSLEs could be used as a diagnostic to validate eddy-resolving oceanic models.

In Fig. 3 (bottom) we see that, for Satellite1/4, the values of FSLEs decay from 0.18 $days^{-1}$ in the southern to 0.03 $days^{-1}$ in the northern area, with similar decays for ROMS1/4. Specifically the North-South difference for Satellite1/4, ROMS1/4 and ROMS1/12 are of the order of 0.15 $days^{-1}$, 0.15 $days^{-1}$ and 0.08 $days^{-1}$, respectively, confirming a lower latitudinal gradient for the case of ROMS1/12. These values do not change much when it is averaged over the 3 degrees stripe offshore (Fig. 3, top), although in this case relative maxima and minima appear, probably in relation with the complex and variable shelf circulation.

The mixing behavior can be also assessed by looking at a proxy of the intensity of 252 mesoscale activity, the Eddy Kinetic Energy (EKE), as done in Gruber et al. (2011). Fig. 4 253 shows that there are regions, as in the FSLE case, with distinct dynamical characteristics. 254 Larger values appear in the south and smaller in the north. This distribution is in good agreement with the one deducted from the FSLEs (Fig. 2). Some simple spatial correla-256 tion (not shown) indicate that EKE and FSLE patterns are well correlated when using a 257 non-linear fitting (power law). For instance, EKE and FSLE computed on the velocity field 258 from Satellite 1/4 exhibit a R^2 of 0.86 for the non-linear fitting: $FSLE = 0.009 \cdot EKE^{0.49}$. 259 It is in agreement with the initial results from Waugh et al. (2006); Waugh and Abraham 260 (2008), for a related dispersion measurement, and confirmed the thorough investigation 261 of the relationship between EKE and FSLE by Hernández-Carrasco et al. (2011b). 262

In the following sections, we study the effect of this variable surface mixing activity on the plankton dynamics.

265 4.2. Plankton dynamics in the Benguela upwelling system.

Evolution of N, P and Z over space and time is obtained by integrating the systems 266 described by Eqs. 8,9,10. The biological model is coupled to the velocity field after the 267 transient time needed to reach stability (60 days). In Fig.5 we show some snapshots of 268 phytoplankton concentrations for the three velocity fields at different times. Since both ROMS simulation were climatologically forced runs, the dates do not correspond to a 270 specific year, whereas we used the actual date for Satellite 1/4. The most relevant feature is 271 the larger value of concentrations near the coast due to the injection of nutrients following 272 Fig. 1. Obviously the spatial distribution of P is dominated by the submeso- and meso-273 scale structures such as filaments and eddies. This is specially noticeable in the south, due to the presence of several Agulhas rings, cyclonic eddies and filaments. Differences 275 are however observed for the three data sets. In particular, it seems that for Satellite 1/4 276 and ROMS1/12 the concentrations extend farther offshore than for ROMS1/4. 277

Several studies (Lehan et al., 2007; d'Ovidio et al., 2009; Calil and Richards, 2010) 278 have shown that chlorophyll distributions in the marine surface are linked to the local 279 maxima or ridges of the FSLEs. This also occurs in our numerical setting, as it is visually 280 shown in Fig. 6. We superimpose contours of high values of FSLE (locating the LCS) 281 on top of phytoplankton concentrations for ROMS1/12 (every 8 days during a 32 days 282 period). In some regions P concentrations are constrained and stirred by lines of FSLE. 283 For instance, the edges of the cyclonic and anti-cyclonic eddies centered at 6 °E, 32 °S, and 28 °S in Fig. 6 on June 11 exhibit large values of phytoplankton concentration. This 285 reflects the fact that tracers, even active such as chlorophyll, still disperse along these 286 LCSs. 287

In order to reveal regions of more intense biological activity, we have computed the temporal average of simulated P. The results, plotted in Fig.7 a), b), c), show that

coastal regions with high P extend approximately, depending on latitude, between half a degree and two degrees offshore. It is comparable with the pattern obtained from the 291 satellite-derived chlorophyll data (Fig. 7 d)). The spatial correlation of averaged simulated 292 chlorophyll with satellite is as follows: $R^2 = 0.85$ for Satellite 1/4 versus Sea WIFS; $R^2 =$ 293 0.89 for ROMS1/4 versus SeaWIFS and $R^2 = 0.85$ for ROMS1/12 versus SeaWIFS. 294 Despite the very simple setting of our models, the phytoplankton development over the 295 Benguela shelf is well simulated by the upwelling parameterization chosen. Note however 296 that our simulated chlorophyll values are about $\simeq 3-4$ times lower than satellite data, 297 as shown by the colorbar scale. Of course several factors, both biological and physical, 298 are not taken into account in this simple setting that might explain this offset. Another 299 possible explanation is the low reliability of the ocean color in very coastal waters optically 300 complex. 301

We now examine the latitudinal distribution of P. The top row in Fig.8 displays the 302 outputs of the numerical simulations that were averaged over a coastal strip of 3° (left) 303 and 6° (right) width. The bottom row is the same but from the satellite chlorophyll data. First of all, phytoplankton biomass has a general tendency to decrease with latitude, an 305 opposite tendency to the ones exhibited by mixing (from FSLEs and EKE) for the three 306 data sets. P values are higher in the northern than in the southern area of Benguela. A 307 common feature is the minimum located just below the Luderitz upwelling cell (28°S), 308 maybe related to the presence of a physical boundary, already studied and named the LUCORC barrier by Shannon et al. (2006) and Lett et al. (2007). Note that on Fig. 3 310 (upper plot), the same latitude was marked by a local maximum of mixing that might 311 be responsible for this barrier. Though not so evident, the same latitudinal tendency is 312 observed for the SeaWIFS data plotted in Fig. 8c) and d). Correlation of zonal average of 313 simulated chlorophyll versus satellite data does not give striking results when considering 314 the whole area (R^2 ranging from 0.1 to 0.5). However, when considering each subsystem 315 independently, high correlation coefficients are found for the south Benguela (R^2 around 316

0.75), but not for the north. It clearly indicates that our simple modelling approach is able 317 tostimulate well the spatial patterns of chlorophyll in the south Benguela, but not properly 318 in the northern part. The 2D vigorous mixing in the south and its associated intense off-319 shore export are sufficient to explain reasonable latitudinal patterns of P. The numerous 320 eddies released from the Agulhas system, moving offshore in the south Benguela, might 321 limit the large development of P by exporting unused nutrients and young phytoplankton 322 communities toward the open ocean, as stated by Gruber et al. (2011). It also suggests 323 that the negative effect seems to be mainly driven by 2D advection toward the open 324 ocean. In the north, other factors seem to play an important role. Among many others, 325 the 3D flow, the shelf width, the rivers and aeolian inputs, the remineralisation pattern, 326 the presence of particular biogeochemical functioning,...etc. have been disregarded from 327 this study, whereas they seem to impact widely plankton dynamics in the north. 328

To address the question of the negative effect of horizontal stirring on phytoplankton 329 concentration in a more quantitative way, we have examined the correlation between 330 these two quantities. We have plotted spatial averages over each subregion (North and 331 South) of every weekly map of FSLE versus the same average of the corresponding weekly 332 map of P, for each week during three years in the case of Satellite and for one year for 333 the case of ROMS (Fig.9). For all cases, a negative correlation between FSLEs and 334 chlorophyll emerges. Thus, the higher the surface stirring/mixing, the lower the biomass 335 concentration. The correlation coefficient is quite similar for all the plots ($R^2=0.80$ to 0.84), and the slopes have the following values: -1 for Satellite1/4, -0.65 for ROMS1/4 337 and -1.5 for ROMS1/12. Note that, similarly to the results of Rossi et al. (2008, 2009) 338 and Gruber et al. (2011), the negative slope is larger but less robust when considering the 339 whole area rather than within every subregion. The suppressive effect of mixing might be 340 dominant only when mixing is intense, as in the south Benguela. Moreover, Gruber et al. 341 (2011) stated that the reduction of biomass due to eddies may extend beyond the regions 342 of the most intense mesoscale activity, not considered here. In fact in our simulations, we 343

observe than averaging FSLEs over a 3° or a 6° coastal band returns quite comparable absolute values, attesting of a significant mixing spreading offshore. However average values of P in Fig. 8 decrease when averaging over a wider area.

The same inverse relationship is observed in Fig.10 using chlorophyll data from SeaWIFS. This analysis confirms the result obtained from satellite velocity fields by Rossi
et al. (2008, 2009) but using FSLEs computed on simulated velocity field with ROMS, at
two different resolutions. In this case, the value of the slopes are: -3.5, -3.4 and -4.7 for
Satellite1/4, ROMS1/4 and ROMS1/12, respectively. The fact that ROMS velocity data
do not necessarily match the dates of SeaWIFS may explain the larger discrepancy in the
values of the correlation coefficient showed in Fig.10.

Then, let us present a brief description of the seasonal behavior of the system. In 354 Fig. 11 we display the temporal evolution over one year of the spatial averages of FSLEs 355 (upper plot) and P (bottom). A climatological average for the case of Satellite1/4 using 356 three years of data. We observe that the seasonal increase in mixing activity (from May 357 to September, roughly winter) is associated to a decrease of the simulated phytoplankton. This also illustrates the seasonal inhibiting effect that the mixing activity has on the 350 phytoplankton dynamics in winter. Note that the seasonal variation of light is not taken 360 into account in our model. However, the temporal variability of plankton in the Benguela 361 is mainly driven by the varying activity of the coastal upwelling cells, reproduced by the 362 function S. 363

Finally, a few sensitivity analysis were done to clarify the role of the 2D advection and
the biological reactions in the simulated plankton fields. For this, we performed virtual
experiments to determine the effect of both processes taken separately. A simulation with
only advection of a passive tracer (without any upwelling parameterization) is compared
to a similar simulation adding the biological reaction terms. The advection-only case
reproduces well the smaller tracer concentrations in the southern domain, whereas the
advection-reaction case presents a more constant latitudinal profile (see Fig. 12). This

confirms that the main influence on the spatial distribution of phytoplankton in the south is 2D advection, with the biological dynamics playing a minor role.

Also, the per capita growth rate of N over time (i.e. $N^{-1}dN/dt$) was computed and 373 averaged over the coastal area in each subsystem to test the mechanisms proposed by 374 Gruber et al. (2011) (see Fig. 13 for the ROMS1/12 simulation). We found that the 375 mean value for each subsystem, North and South, are $-3 \cdot 10^{-5}$ and $-1 \cdot 10^{-4}$ day^{-1} , 376 respectively. This confirms that nutrients are being lost toward the open ocean by simple 377 2D advection almost four times more in the south than in the north. It has to be compared 378 with the mixing activity being about three times higher in the south than in the north 379 (Fig. 3). The same behavior is also observed in the other two cases ROMS1/4 and 380 Satellite (not shown). Note also that the loss of nutrient appear to be maximal in the 381 winter months (maximum mixing), although there is a slight decay in between the two 382 subsystems. 383

5. Conclusions

This study is based on numerical analysis from a simple biological NPZ model coupled 385 with different velocity fields (satellite and model) over the Benguela area. Although in a 386 simple framework, a reduction of phytoplankton concentrations in the coastal upwelling 387 for increasing mesoscale activity has been successfully simulated. Horizontal stirring was 388 estimated by computing the FSLEs and was correlated negatively with chlorophyll stocks. Similar results are found, though not presented in this manuscript, for the primary pro-390 duction, defined as the first term in F_P (Eq.5), i.e. $PP = \beta \frac{N}{\kappa_N + N} P$. Some recent 391 observational and modelling studies proposed the "nutrient leakage" as a mechanism to 392 explain this negative correlation. Here we argue that Lagrangian Coherent Structures, 393 mainly mesoscale eddies and filaments, transport a significant fraction of the recently 394 upwelled nutrients nearshore toward the open ocean before being efficiently used by the 395 pelagic food web. Although some studies dealt with 3D effect, we have shown that 2D 396

advection processes seems to play an important role in this suppressive effect. Our analvsis suggest that the inhibiting effect of the mesoscale activity on the plankton occurs 398 when the mixing reach high levels, as in the south Benguela. However, this effect is not 399 dominant under certain levels of turbulence. We have also shown that the inhibiting effect 400 of intense mixing is maximal during the winter months. It might indicate that planktonic 401 ecosystems in oceanic regions with vigorous mesoscale dynamics can be, as a first ap-402 proximation, easily modeled just by including a realistic flow field. The small residence 403 times of waters in the productive area will smooth out all the other neglected biological 404 factors in interaction. However, these factors are required when modelling an oceanic 405 regions with low mixing, associated with high residence time leading to the predominance of complex combinations of factors.

Our findings confirm the unexpected role that mesoscale activity has on biogeochemical dynamics in the productive coastal upwelling. Strong vertical velocities are known to be associated with these physical structures and they might have another direct effect by transporting downward rich nutrient waters below the euphotic zone. Further studies are needed such as 3D realistic modelling that take into account the strong vertical dynamics in upwelling regions to test the complete mechanisms involved.

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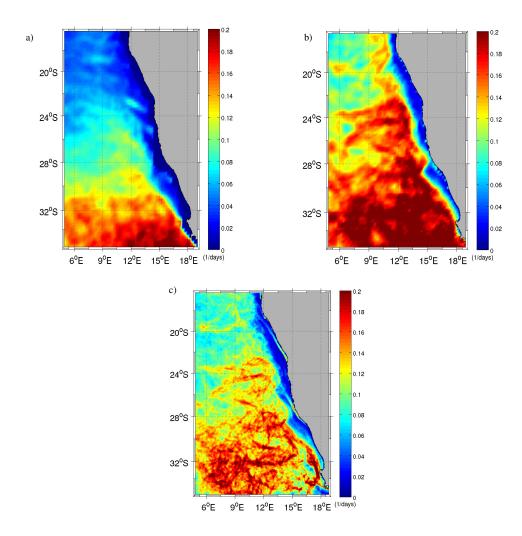
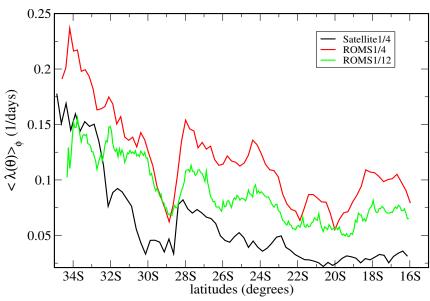


Figure 2: Spatial distribution of time average of weekly FSLE maps in the Benguela region. a) Three years average using data set Satellite1/4; b) one year average using ROMS1/4; c) one year average using ROMS1/12. The units of the colorbar are 1/days.

Average over 3 degrees offshore



Average over 6 degrees offshore

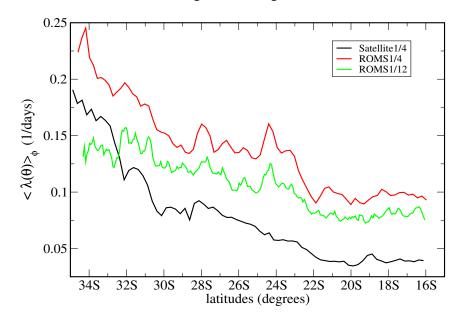


Figure 3: Zonal average over coastal bands of the FSLE time averages from Fig. 2 as a function of latitude. Top) From the coast to 3 degrees offshore; bottom) to 6 degrees offshore.

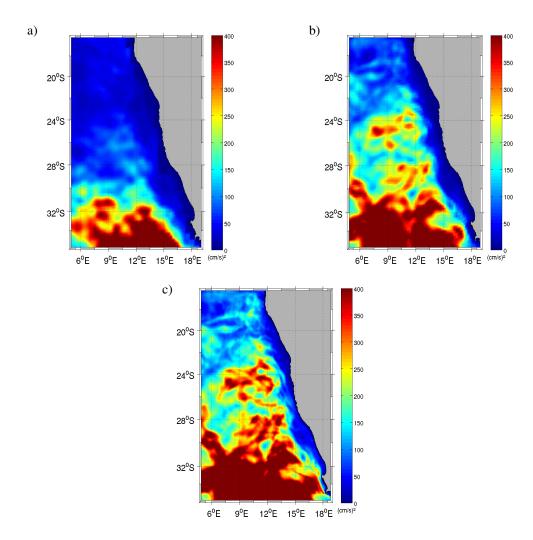


Figure 4: Spatial distribution of annual EKE in the Benguela region. a) using velocity data from Satellite at spatial resolution $1/4^{\circ}$ (Satellite1/4) b) using velocity data from ROMS at spatial resolution $1/4^{\circ}$ (ROMS1/4) c) using velocity data from ROMS at spatial resolution $1/12^{\circ}$ (ROMS1/12). The units of the colorbar are $(cm/s)^2$

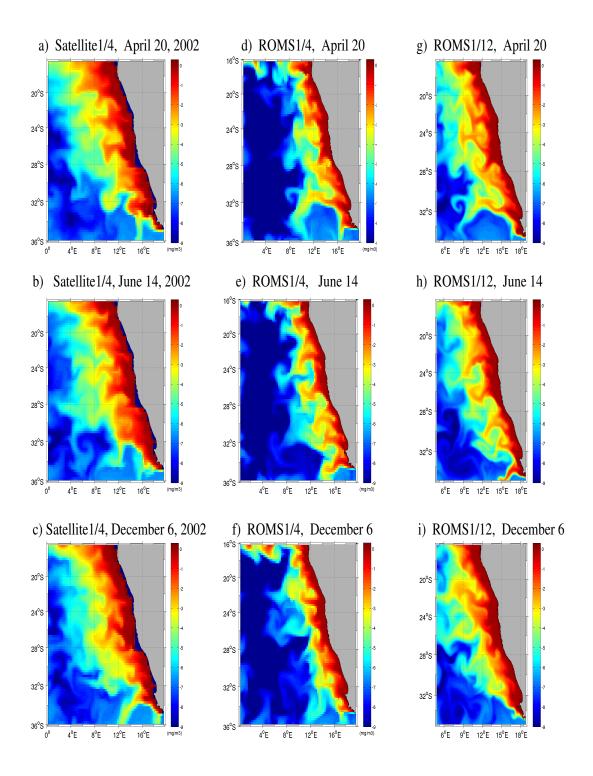


Figure 5: Snapshots of spatial distribution of phytoplankton concentration from the simulations: Left column) corresponding to the simulation using Satellite1/4; Middle column) ROMS1/4; Right column) from ROMS1/12. Logarithmic scale is used to improve the visualization of the structures. The units for the colorbar are mg/m^3

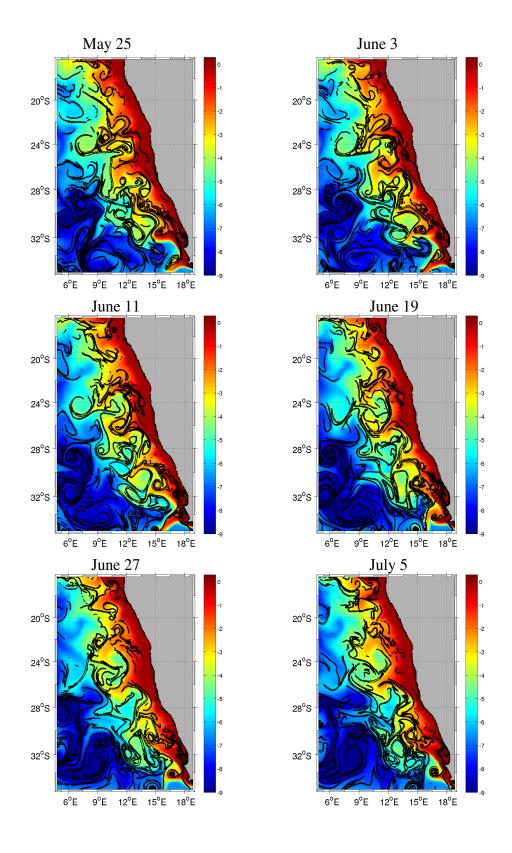


Figure 6: Snapshots every 8 days of large (top 30%) values of FSLE superimposed on P concentrations calculated from ROMS1/12 in mg/m^3 . Logarithmic scale for phytoplankton concentrations is used to improve the visualization of the structures

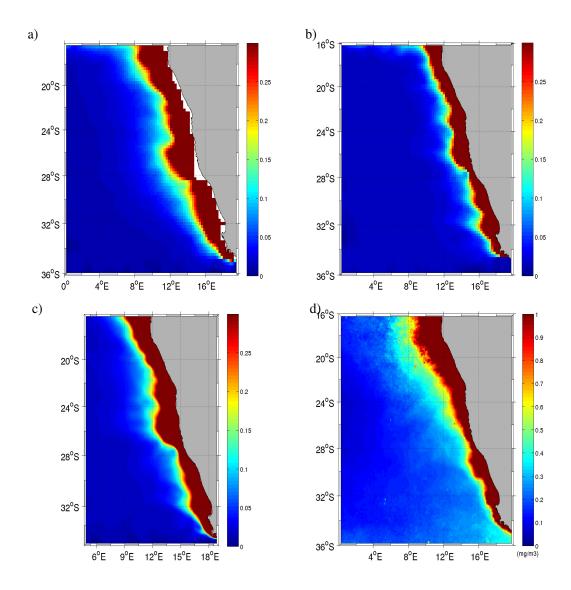


Figure 7: Spatial distribution of the time average of phytoplankton concentrations: a) Three years average using Satellite1/4, b) One year average from ROMS1/4, c) One year average from ROMS1/12, d) Three years average of monthly SeaWIFS data. The units of the colorbar are mg/m^3 .

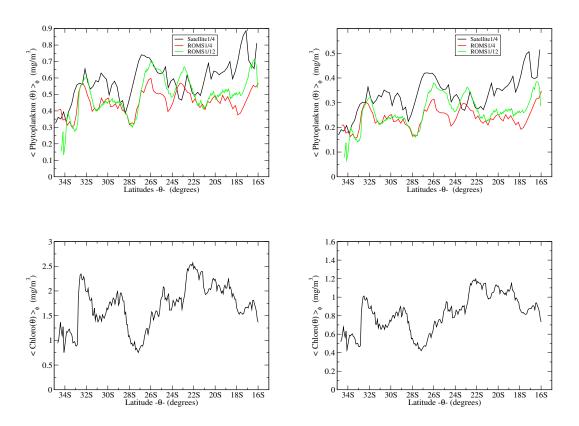


Figure 8: Zonal mean, over a 3 degrees (left) and 6 degrees (right) width coastal band, of the time averages of modelled phytoplankton (upper plots) and derived from satellite (lower plots) plotted as a function of latitude.

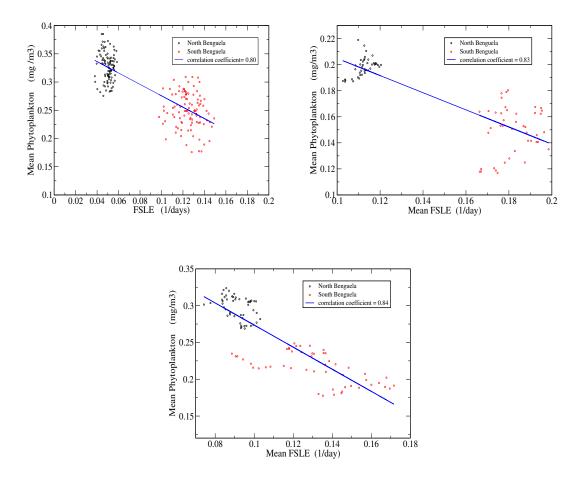


Figure 9: Weekly values of spatial averages of phytoplankton versus weekly values of spatial averages of FSLE, where the average are over the North and South subareas of Benguela. a) Satellite1/4, b) ROMS1/4 and c) ROMS1/12

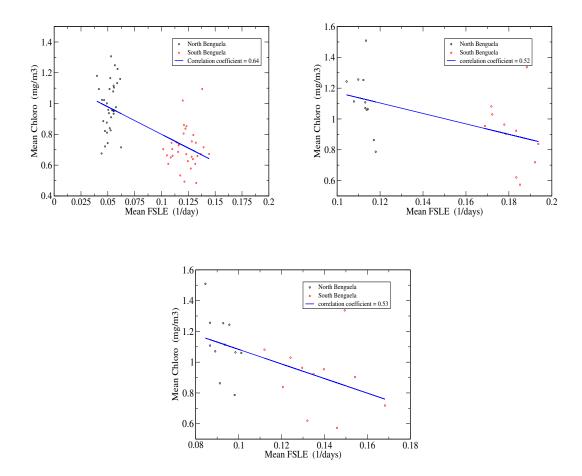
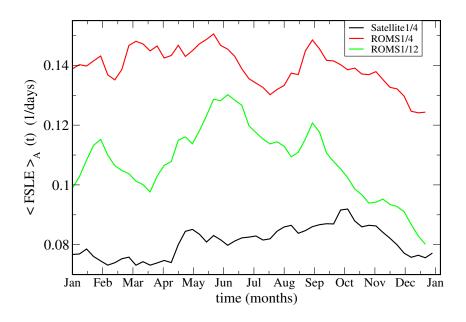


Figure 10: Monthly values of spatial averages of Chlorophyll from SeaWIFS data versus spatial average of FSLE, where the average are over the North and South subareas of Benguela. FSLE values are from a) Satellite1/4, b) ROMS1/4 and c) ROMS1/12.



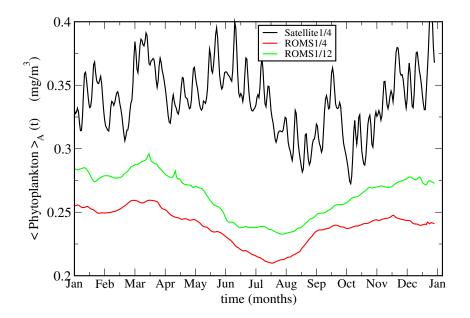


Figure 11: Temporal evolution of horizontal mixing (Spatial average of FSLEs) for the three velocity data sets (top). Temporal evolution of spatial averages of simulated phytoplankton for the three velocity data sets (bottom).

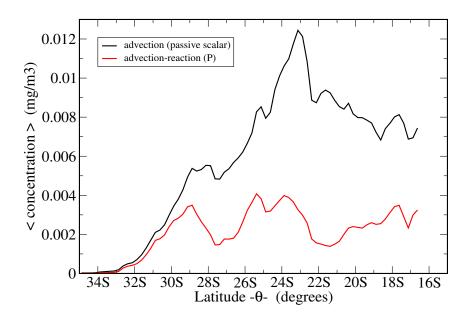


Figure 12: Normalised comparison of the time averages of a passive scalar (advection only) and of P (advection-reaction), as a function of latitude.

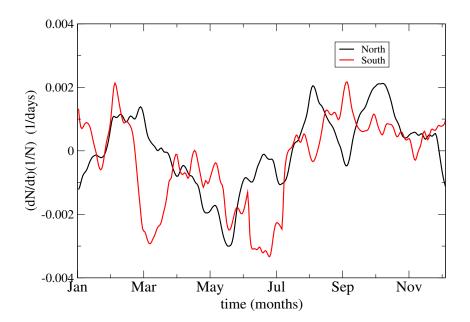


Figure 13: Time evolution of the spatial average of the per capita growth rate of nutrients for the ROMS1/12 case.